

[REPLACEMENT SHEET]  
Application No. 10/516,361

FPAA/288(PCT)

5'-GTT TTT GTG GTA GTA TGT GAT TTA GTC ATT CAA CCC TCG TAG TGG GCG CAA ACG CTC TAA CTT AAA TGT A-J'

(SEQ ID No: 32)

Fig 8

[REPLACEMENT SHEET]  
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FPAA/288(PCT)

5'-TGC GGG GTA CTA CAG CGC CCT GAC CAT GGC CAT CTT CCA GGA CCT CGG-3'

(SEQ ID No: 33)

Fig 9

[REPLACEMENT SHEET]  
Application No. 10/516,361

FPAA/288 (PCT)

5'-ACG GAG CGG CTG AAG GTG CGG CAG GTG CAG GAC AAG TGG A-3'

(SEQ ID No: 34)

**Fig 10**

[REPLACEMENT SHEET]  
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FPAA/288 (PCT)

**5'-ATG GCG CCT GCC TCG GAT GCG GGG TAC TAC AGC GCC-3'**

(SEQ ID No: 35)

**Fig 11**

[REPLACEMENT SHEET]  
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FPAA/288 (PCT)

**5'-ACT TAA GTT AGA GCG TTT GC-3'** (SEQ ID No: 1)

|  
**FAM**

**Fig 12**

[REPLACEMENT SHEET]  
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FPAA/288(PCT)

5'-GGG GTA CTA CAG CGC CCT GA-3' (SEQ ID NO: 10)  
|  
FAM

Fig 13

FPAA/288(PCT)

5' - GTC CTG GAA GAT GGC GAT GG-3' (SEQ ID NO: 13)

|  
JOE

Fig 14.

[REPLACEMENT SHEET]  
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FPA&/288(PCT)

5' - GGG GTA CTA CAG CGC CCT-3' (SEQ ID No: 21)  
|  
FAM

Fig 15

[REPLACEMENT SHEET]  
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FPAA/288 (PCT)

DABCYL-5'-ATG GCC ATC GTC CTG GAA GAT GGC CAT GG-3'

(SEQ ID No: 23)

JOE

Fig 16

[REPLACEMENT SHEET]  
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FPAA/288(PCT)

5'-DABCYL-ATG GCC ATC GTC CTG GAA GAT GGC CAT GG-3'

| (SEQ ID No:22)  
FAM

Fig 17

[REPLACEMENT SHEET]  
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FPAA/288(PCT)

5' -GCT CAT GGC GCC TGC CTC G-3' (SEQ ID NO: 24)  
|  
DABCYL

Fig 18

[REPLACEMENT SHEET]  
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FPAA/288(PCT)

5'-TGCACACCGA GCGGCTGAAG GTGCGGCAGG TGCAGGACAA  
GTGGAAGGTG ACGGGCATGG GCAACGAGAT CTGTGGCCAC  
TTCAAGGTGC CGCCGGCGCA CATCACCGAT GGCCTGAGCA  
ACACCGACTT CGTGATGTAC GTGCCTCCG TGCCGAGCGA  
GGGGGATGTG CTGGCGTGGG CCACGACCTG CCAGGTGTT  
TCTGACGGCC ATCCAGCCGT GGGCGTCATC AACATCCCCG  
CGGCGAACAT TGCCTCGCGG TACGACCAGC TGGTGACGCG  
TGTCTCACG CACGAGATGG CGCACCGCGCT CGGCTTCAGC  
GTCGTCTTCT TCCGAGACCG CCGCATCCTG GAGAGCATT  
CGAACGTTCG GCACAAGGAC TTGATGTTT CCGTGATCAA  
CAGCAGCACG GCGGTGGCGA AGGCGCGCGA GCAGTACGGC  
TGCAGCACCT TGGAGTATCT GGAGATGGAG GACCAGGGCG  
GTGCGGGCTC CGCCGGGTCTG CACATCAAGA TGCAGCACGC  
GCAGGACGAG CTCATGGCAC CTGCCTCGGA TGCAGGGTAC  
TACAGCGCCC TGACCATGGC CATCTTCCAG GACCTCGGCT  
TCTACCAGGC; 3' (SEQ ID NO: 36)

Fig 19